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The invention in the wood of the control of pupplies have been controlled been inventional and a per a trade and a per a
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misma immundeficienty vitus 1; HIV-1; endotoxaemia; septic shock; 
preservative; sterilant; food additive; topical agent.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Novel isolated cationic, polyphemusin-like peptide having two mati-parallel beta strands stabilized by two disulfible bonds, beta hairpin loop and mutimicrobial activity useful to linkibit bacterial, hairpin loop and mutimicrobial activity useful to
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heelyd78; i.imilus polyphemusin
(from "kam872.pep")
TOIG of: aael9478 check: 3063 from: 1
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manna immunodeficatery vitus I; HTV-1; endocrosemia; septic shock;
preservative; sterilant; food additive; topical agent
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Indirect file
Sequence or key file
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Hit display
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action 1 match found in sequence:

(from "kam872.pep")

TOIG of: nee19477 check: 3126 from: 1 to: 18
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Thu Aug 28 07:33:11 2003
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Nucleic acid code matching
Find non-matching hits only
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Display full annotations
Sequence context
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The invention that was optimized, polyphematical periods being a managed and a beta content of the period of the invention are useful (of inhibiting the propert of increase such as person pointive (e.g., Staphylococcas ances, propert of increase such as temp positive (e.g., Staphylococcas ances, period of increase and a period of incre
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                                                                                                                                                                                                          (UYBR-) UNIV BRITISH COLUMBIA.
                                                               27-JUN-2001; 2001WO-CA00918.
                                                                                                                                     27-JUN-2000; 2000US-0604864.
                                                                                                                                                                                                                                                                                        Bancock REW, Shang L;
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03-JAN-2002.
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AAE19478 Length: 18 August 28, 2003 07:00 Type: P Check: 3063 Pound using 'seq17' (kam872.key) Sequence 18 AA;

RVWCRYRCYRGFCRRFCR

-- Search Statistics --Times:

Total Elapsed 00:00:00.00

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Number of sequences searched: Number of sequence hits: Number of separate matches: Number of sequence hits saved:

! FINDPATTERNS on PIR: * allowing 0 mismatches

Thu Aug 28 07:32:52 2003

1 WCF(F,Y,W,A,L,I,V)VC(K,R,H,F,Y,W)RG(K,R,H)CR(K,R,H,F,Y,W)KCRR

Databases searched: NBRF, Release 76.1, Released on 12May2003, Formatted on 10Jun2003

0 96,168,682 283,308 02:27.89 Total finds: Total length: Total sequences: CPU time:

Thu Aug 28 07:32:54 2003

1 FINDPATTERNS on Swiss-Prot:* allowing 0 mismatches

1 WCF(F,Y,W,A,L,I,V)VC(K,R,H,F,Y,W)RG(K,R,H)CR(K,R,H,F,Y,W)KCRR

Databases searched: SWISS-PROT, Release 41.1, Released on 6Jun2003, Pormatted on 9Jun2003

47,026,705 127,863 01:14.46 Total finds: Total length: Total sequences: CPU time:

1 WCF(F,Y,W,A,L,I,V)VC(K,R,H,F,Y,W)RG(K,R,H)CR(K,R,H,F,Y,W)KCRR

Databases searched:
STYLENBU, Release 23.0, Released on 49ar2003, Formatted on Thar2003
TOTAL Lines 128, 052, 60
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human immunodeficiency virus 1; Priv. 1; endovarseis, septic shock;
preservative; sterilant; food additive; topical agent.
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aac19476; Limius polyphemus polyphemusin-like peptide #7.
(from 'kam872.pep')
TOIS of: aac19476 oheck: 4644 from: 1 to: 19
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mamniamundeficiency virus 1; HIV-t); endetrosemia; septic shock;
preservative; sterilant; food additive; topical sqeit.
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                                                                                                                                                                             Selected search type is key against sequence data banks or files. Selected scope is Sequence respectively. Representations key from 'Ram972.key's segif (AA) ID seqif AA preliminary pattern
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Sequence or key file
List of hits
Hit display
Name and annotations
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                                                                  Quest - Quick User-directed Expression Search Tool Release 5.4
                                                                                                                                                -- Outline of search "seq16ags"
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Find non-matching hits only
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followed by
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the invention of solution to contain the properties bring an attachment of the contain the contain of periods but the contain of periods of the contain o
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 3; Page 35; 57pp; English.
                                                                                                                                                                                                             (UYBR-) UNIV BRITISH COLUMBIA.
                                                                    27-JUN-2001; 2001WO-CA00918,
                                                                                                                                 27-JUN-2000; 2000US-0604864.
                                                                                                                                                                                                                                                                                 Bancock REW, Zhang L:
                                                                                                                                                                                                                                                                                                                                                     WPI; 2002-226915/28
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03-JAN-2002.
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RENCFIVCRRGRCYVACRR

-- Search Statistics --

CPU 00:00:00 Number of sequences searched: Number of sequence hits: Number of separate matches: Number of sequence hits saved: Times:

Total Elapsed 00:00:00.00 7000 seq15pir.find

| FINDPATTERNS on PIR: * allowing 0 mismatches

Thu Aug 28 07:32:54 2003

1 (K,R,H,F,Y,W)RWCPRVCY(K,R,H)G(K,R,H,P,Y,W)C(K,R,H)(F,Y,W,A,L,I,V)(K,R,H,F,Y,W

Databases searched: NBRF, Release 76.1, Released on 12May2003, Formatted on 10Jun2003 0 96,168,682 283,308 05:31.47 Total finds: Total length: Total sequences: CPU time:

! FINDPATTERNS on Swiss-Prot: * allowing 0 mismatches

1 (K,R,H,F,Y,W)RWCFRVCY(K,R,H)G(K,R,H,F,Y,W)C(K,R,H)(F,Y,W,A,L,I,V)(K,R,H,F,Y,W

Databases searched: SWISS-PROT, Release 41.1, Released on 6Jun2003, Pormatted on 9Jun2003

47,026,705 127,863 02:43,51 Total finds: Total length: Total sequences: CPU time:

1 (K,R,H,F,Y,W)KWCPRVCY(K,R,H)G(K,R,H,P,Y,W)C(K,R,H)(F,Y,W,A,L,I,V)(K,R,H,F,Y,W ! FINDPATTERNS on sptrembl:* allowing 0 mismatches

Databases searched: SPTREMEL, Release 23.0, Released on 4Mar2003, Formatted on 7Mar2003 Total finds: 258,052,604 Total sequences: 830,525 Total sequences: 15:20,59 CPU time: 15:20.39

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The invention of entires to contract, gaptymentain; like peticles having the invention of the invention of the invention of the atreads subthitsed by you can set distribute books and a beta bright along the peticles of the invention are useful for inhibiting the profit of standards such as garden positive (e.g., Simphiconess arms, profit of standards such as garden positive (e.g., Simphiconess arms, profit of standards and present in the standards and security in the standards of the standards are contracted in the standards are contracted in the standards are standards as the standards are standards as the standards are standards as the standards are contracted indextor (e.g., Septic should) as present associated indextor (e.g., Septic should) as presentation, as bread spectrum antistarchial spent (diserior theory) as present associated indextor (e.g., Septic should) as present as associated indextor (e.g., Septic should) as present as presental and spectrum antistarchial spent (diserior toward) and interest of particular standards are of periodic as present as a contract sequence in a capture of the special spent (Septic Septic Se
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                                                                                                                                                                                                                                                                    Novel isolated cationic, polyphemusin-like peptide having two
anti-parallel beta strands stabilised by two disulfide bonds, beta
hairphi loop and antimicrobial activity useful to inhibit bacterial,
(yast, vital groth)
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assla473 : Limilus polyphemus polyphemusin-like peptide #4.
(from *kam872.pop*)
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1 8. .13
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                                                                                                                                              (UYBR-) UNIV BRITISH COLUMBIA
                                                              27-JUN-2001; 2001WO-CA00918.
                                                                                                        27-JUN-2000; 2000US-0604864.
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                                                                                                                                                                                            Zhang L;
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Disulfide-bond 8
Modified-site 1
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                                                                                                                                                                                            Bancock REW,
                    03-JAN-2002.
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242422222222222
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hamm.lamminoficitency virus 1; RTV-1; andocrosemia; septic shock;
preservatives sterilant; food additive; topical agent.
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Selected sequence. Key from "kam27.key":
Seq15 (Ah] ID seq15 Ah preliminary pattern
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Sequence or key file
List of hits
Hit display
Name and annotations
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see19472 ; Janualus polyphemusin-11ke peptide #3.
(from *ram32.pep*)
FOUG Of: ase19472 check: 3228 from: 1 to: 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Limulus polyphemus polyphemusin-like peptide #3.
                                                                                                                                                  -- Outline of search "seql5ags" --
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                                                                               Quest - Quick User-directed Expression Search Tool
Release 5.4
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Nucleic acid code matching
Find non-matching hits only
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Notify at end of run
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Display full annotations
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                            /note- "Amidated"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 3; Page 35; 57pp; English.
                                                                                                                                                                                                                                                                                                                       (UYBR-) UNIV BRITISH COLUMBIA,
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Number of sequence hits:
Number of separate matches:
Number of sequence hits saved:
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18
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Modified-site
                                                                                    W0200200687-A2
                                                                                                                                             03-JAN-2002.
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     The invention shalts to catchin, polyphemalities periods being the attended being the beat strong stabilised by two more dishilate bond and a beta brains stabilised by two more dishilate bond and a beta brains include so that the properties of the invention are safeth for inhibiting the proph of attended such as impositive (e.g., Suphyloconous access, proch of attended such as impositive (e.g., Suphyloconous access, proch of attended such as impositive (e.g., Suphyloconous access, proch of attended such as impositive (e.g., Suphyloconous access, and the suppositive (e.g., Suphyloconous access) of the suppositive to decrease (e.g., Suphyloconous access (e.g., Suphyloconous access) of the suppositive to decrease (e.g., Suphyloconous access (e.g., Suppositive (e.g., Suppositive) of the suppositive (e.g., Suppositive) 
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mann immunodeficiency vitus 1; HIV-1; endetozemás, septic shock;
preservatives, tetrilan; food additive; topical agent.
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Novel isolated cationic, polyphemusin-like peptide having two anti-parallel beta strands stabilized by two disulfide bonds, beta hairpin loop and antimitrobial activity useful to inhibit bacterial, yeast, vital growth.
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aaei19474 ; Limulia polyphemus polyphemusin-like peptide #5.
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/note- "Amidated"
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                                                        WO200200687-32
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Total Elapsed deco

! 1 REMCP(F,Y,W,A,L,I,V)VC(K,R,H)RG(R,V,A)CY(R,V,A)(R,V,A)CR

Databases searched: NBRF, Release 76.1, Released on 12May2003, Formatted on 10Jun2003

96,168,682 283,308 02:31.79 Total finds: Total length: Total sequences: CPU time:

Thu Aug 28 07:32:55 2003

! FINDPATTERNS on Swiss-Prot: * allowing 0 mismatches

1 RRWCF(F,Y,W,A,L,I,V)VC(K,R,H)RG(R,V,A)CY(R,V,A)(R,V,A)CR

Databases searched: SWISS-PROT, Release 41.1, Released on 6Jun2003, Formatted on 9Jun2003

47,026,705 127,863 01:14.99 Total finds: Total length: Total sequences: CPU time:

! FINDPATTERNS on sptrembl: * allowing 0 mismatches

1 RENCE(F,Y,W,A,L,I,V)VC(K,R,H)RG(R,V,A)CY(R,V,A)(R,V,A)CR

Databases searched: SPTREMBL, Release 23.0, Released on 4Mar2003, Formatted on 7Mar2003

258,052,604 830,525 07:14.40 Total finds: Total length: Total sequences: CPU time:

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the invention obtains to statistically, plyphasmin-like postides bring a minimizational antityty. These cutionic species have to earlightnish the statistic statistic bods and a beta bring statistic statistic bods and a beta bring in close periods of the invention are useful for inhibiting the provide of attaches such as stan pointive (e.g., Suphylococcus ancess, provide of attaches such as stan pointive (e.g., Suphylococcus ancess, c.g., Supply concess ancess, provide of attaches such as paraporative (e.g., Supply concess ancess, c.g., Supply concess ancess, such as a supply of the s
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to Ail odour producing microbes (Micrococci). The present sequence is
Limitus polyphemus polyphemus polyphemus in-like peptide.
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minan immunodeficiency virta i, HTV-1 endotoxamia; septic shock;
preservative; sterilant; food additive; topical agen:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Novel isolated cationic, polyphemusin-like peptide having two marti-parallel beta strands tabilitzed by two disulfide bonds, beta hamipin loop and autimicrobial activity useful to lambit bacterial,
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aae19471 ; Limulus polyphemus polyphemusin-like peptide #2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 3; Page 35; 57pp; English.
                                                                                                                                                                                                                                                                                                              (UYBR-) UNIV BRITISH COLUMBIA
                                                                                                                                                                                27-JUN-2001; 2001WO-CA00918.
                                                                                                                                                                                                                                               27-JUN-2000: 2000US-0604864.
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                                                                                                                                                                                                                                                                                                                                                                                   Zhang L;
                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2002-226915/28.
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TOIG of: aae19471
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      seq14ags.res
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himman immunodeficiency rruns i; mirty-i endocamenta; septic shock;
preservative; sterilant; food additive; topical agent.
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aae19470; illmulus polyphemus polyphemusin-like peptide #1.
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                                                                                                                                                                                                                                                                                     Quest - Quick User-directed Expression Search Tool
Release 5.4
                                                                                                                                                                                                                                                                                                                                                                                                      -- Outline of search "seq14ags"
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Selected sequence key from "kam872.key":
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Notify at end of run
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hard-parallel beta strands stabilited by two distified bonds, beta
haitpin loop and antimicrobial activity useful to inhibit bacterial,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 3; Page 35; 57pp; English.
                                                                                                                                                                             (UYBR-) UNIV BRITISH COLUMBIA,
                                         27-JUN-2001; 2001WO-CA00918.
                                                                                                       27-JUN-2000; 2000US-0604864.
                                                                                                                                                                                                                                                  Hancock REW, Zhang L;
                                                                                                                                                                                                                                                                                                                       WPI; 2002-226915/28.
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WCFAVCYRGRCRRKCRR

CPU 00:00:00 Times:

Total Elapsed 00:00:00.00

-- Search Statistics --

Number of sequences searched: Number of sequence hits: Number of sequence hits saved: Number of sequence hits saved:

I R(F,Y,W,A,L,I,V)WC(K,R,H)(K,R,H,F,Y,W)(K,R,H)CYRGFC(K,R,H)(K,R,H,F,Y,W)(F,Y,W ! FINDPATTERNS on PIR:* allowing 0 mismatches

Databases searched: NBRF, Release 76.1, Released on 12May2003, Formatted on 10Jun2003 Total finds: Total length: Total sequences: CPU time:

96,168,682 283,308 02:52.77

Thu Aug 28 07:32:56 2003

! FINDPALTERNS on Swiss-Prot:* allowing 0 mismatches

1 R(F,Y,W,A,L,I,V)NC(K,R,H)(K,R,H,F,Y,W)(K,R,H)CYRGFC(K,R,H)(K,R,H,F,Y,W)(F,Y,W

Databases searched: SWISS-PROT, Release 41.1, Released on 6Jun2003, Formatted on 9Jun2003

47,026,705 127,863 01:26.17 Total finds: Total length: Total sequences: CPU time:

1 FINDPATTERNS on sptrembl:* allowing 0 mismatches

1 R(F,Y,W,A,L,I,V)WC(K,R,H)(K,R,H,F,Y,W)(K,R,H)CYRGFC(K,R,H)(K,R,H,F,Y,W)(F,Y,W

Databases searched: SPTREMBL, Release 23.0, Released on 4Mar2003, Formatted on 7Mar2003

Total finds: Total length: Total sequences: CPU time:

258,052,604 830,525 08:15.75

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haitpin loop and antimicrobial activity useful to Inhibit bacterial,
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                                                                                                                                                                                                                                                                                                                               Claim 3; Page 35; 57pp; English.
(UYBR-) UNIV BRITISH COLUMBIA.
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Number of separate matches:
Mumber of sequence hits saved
                                                              Bancock RBW, Shang L;
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                                                                                                                       WPI; 2002-226915/28
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preservative; sterilant; food additive; topical, agent.
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No
Yes
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Selected sequence Key from "kan072.key":
seq18 (AA) ID seq18 AA preliminary pattern
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Sequence or key file
List of hits
Hit display
Name and annotations
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each 77; Inmils polyphemus polyphemusin-like peptide #10.
(form *ksm872.pep') check: 3049 from: 1 to: 18
701G of: aae19479 check: 3049 from: 1 to: 18
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                                                                                                                       Quest - Quick User-directed Expression Search Tool
Release 5.4
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Nucleic acid code matching
Find non-matching hits only
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Fime to start comparison
Notify at end of run
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Note position of hit
Display full annotations
Sequence context
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  File : kam872.pep
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Limulus polyphemus
> 0 < 0 | 0 IntelliGenetics > 0 <
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                                                                                                                                                                                                                                                                                                                                                           seq18 (AA) ID se
followed by
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Selected files:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAE19479;
                                                                                                                                                                                                                                                                                                                                                                                                                          TEMO
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August 27

! FINDPATTERNS on PIR:* allowing 0 mismatches

1 REWC(K, R, H) RVCY(R, Y, W, A, L, I, V) GECYRKCR

Databases searched: NBRF, Release 76.1, Released on 12May2003, Formatted on 10Jun2003

Total finds: Total length: Total sequences: CPU time:

0 96,168,682 283,308 02:08.60

seq18sp.find

August 27 | FINDPATTERNS on Swiss-Prot: * allowing 0 mismatches 1 RRWC(K,R,H)RVCY(P,Y,W,A,L,I,V)GPCYRKCR

Databases searched: SRISS-TRO7, Melease 41.1, Meleased on Glun2003, Formatted on SJun2003

47,026,705 127,863 01:05.55 Total finds: Total length: Total sequences: CPU time:

Thu Aug 28 07:32:56 2003

August 27

Databases searched: SPTREMBL, Release 23.0, Released on 4Mar2003, Formatted on 7Mar2003

258,052,604 830,525 06:07.82

Total finds: Total length: Total sequences: CPU time:

! FINDPATTERNS on sptrembl:* allowing 0 mismatches 1 REWC(K, R, H) RVCY(F, Y, W, A, L, I, V) GFCYRKCR

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The inventor incluses to catching peptides have been inventor that the catching of the inventor inclusion to catching the catching of the catc
                                                                                                                                  Novel isolated cationic, polyphemusin-like peptide having two anti-parallel beta strands stabilized by two disulfide bonds. Deta halrphi loop and antimicrobial activity useful to inhibit bacterial, yeast, vital growth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAR19480 Length: 19 August 28, 2003 07:00 Type: P Check: 4675 ound using 'seq19' (kam872.key)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Total Elapsed
00:00:00.00
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                -- Search Statistics --
                                                                                                                                                                                                                                                      Claim 3; Page 35; 57pp; English.
(UYBR-) UNIV BRITISH COLUMBIA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               00:00:00:00
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  sequence hits:
separate matches:
sequence hits saved:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            sequences searched:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         REWCFRVCYRGRFCYRKCR
                                            Hancock REW, 2hang L;
                                                                                         WPI: 2002-226915/28.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 19 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Found using 'seq19'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Number of s
Number of s
Number of s
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Times:
    Polyphemusin-like peptide; antimicrobial; sepsis-associated disorder;
human immunodeficiency virus 1; HTV1; endocksemia, septic shock;
preservative; steatlant; food additive; topical agent.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            No Kes
                                                                                                                                                                                                Selected search type is key against sequence data banks or files, selected sequence key from 'kan872.key': seq19 (AA) ID Se437 AA PREJABINARY pattern tollowed by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        match found in sequence:
ase19480 ; Limulus polyphemus polyphemusin-like peptide #11.
(from *kam872.pep' 72.pep' 7503 of from: 1 to: 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      File Options:
Indirect file
Sequence or key file
List of hits
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Hit display
Name and annotations
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Limulus polyphemus polyphemusin-like peptide #11.
                                                                            Quest - Quick User-directed Expression Search Tool
Release 5.4
                                                                                                                                                        -- Outline of search "seq19ags"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         -- Output Parameters --
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                -- Run Parameters --
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Location/Oualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAE19480 standard; peptide; 19 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /note- "Amidated"
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now
No
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No
Yes
Yes
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  31-MAY-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Nucleic acid code matching
Find non-matching hits only
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Run mode
Time to start comparison
Notify at end of run
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Report key used
Note position of hit
Display full annotations
Sequence context
                                                                                                                                                                                                                                                                                                                                                                                                                                                            File : kam872.pep
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k or r or h
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> 0 < Olimination of to IntelliGenetics
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Disulfide-bond
Modified-site
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                                                                                                                                                                                                                                                                                                                                                                                                               Selected files:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          format Options:
                                            0
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August 27, 2003 06:54

! FINDPATIERNS on PIR:* allowing 0 mismatches

Databases searched: NBRF, Release 76.1, Released on 12May2003, Formatted on 10Jun2003 1 REWCFRVCYRG(K,R,H)FCYRKCR

96,168,682 283,308 01:57.78

Total finds: Total length: Total sequences: CPU time:

1 FINDPATTERNS on Swiss-Prot: * allowing 0 mismatches Thu Aug 28 07:32:57 2003 1 RRWCFRVCYRG(K,R,H)FCYRKCR Databases searched: SRISS-PROT, Release 41.1, Released on GJun2003, Pormatted on 9Jun2003

47,026,705 127,863 59.41

Total finds: Total length: Total sequences: CPU time:

August 27, 2003 06:55

August 27, 2003 06:48

: FINDPATTERNS on sptrembl:* allowing 0 mismatches

Databases searched: SPTRBMEL, Release 23.0, Released on 4Mar2003, Formatted on 7Mar2003

1 RRWCFRVCYRG(K, R, H)FCYRKCR

Total finds: Total length: Total sequences: CPU time: